

**Version With Markings to Show Changes Made**

In the specification page 11, lines 7 through 15 have been amended as follows:

The PCR primers used were: BMP9, forward primer: 5'-TCCCCACCGACTTGTTCTTC-3' (SEQ ID NO: 1), reverse primer: 5'-GAGAGTCAGCTGGGAGCTTGA-3' (SEQ ID NO: 2). GAPDH, forward primer: 5'-TGTGTCCGTCGTGGATCTGA-3' (SEQ ID NO: 3), reverse primer: 5'-CCTGCTTCACCACTTCTTGA-3' (SEQ ID NO: 4). RT-PCR for the M exon of ChAT was performed using the Access RT-PCR system from Promega. The forward primer (5'- GGG GTG GCT GGT TTG CTT GCA GTC A -3') (SEQ ID NO: 5) was designed specifically for detection of transcripts originating at the M promoter, and the reverse primer (5'- GGG GGC ACT GGC AAC TTA GGT AAG -3') (SEQ ID NO: 6) was derived from the coding region of the ChAT gene.